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APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO.
10/505,239	10/12/2004	Nadya I Tarasova	229694	1908
45733	7590	12/28/2007	EXAMINER	
LEYDIG, VOIT & MAYER, LTD. TWO PRUDENTIAL PLAZA, SUITE 4900 180 NORTH STETSON AVENUE CHICAGO, IL 60601-6731			KHANNA, HEMANT	
		ART UNIT	PAPER NUMBER	
		1654		
		MAIL DATE	DELIVERY MODE	
		12/28/2007	PAPER	

Please find below and/or attached an Office communication concerning this application or proceeding.

The time period for reply, if any, is set in the attached communication.



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APPLICATION NO./ CONTROL NO.	FILING DATE	FIRST NAMED INVENTOR / PATENT IN REEXAMINATION	ATTORNEY DOCKET NO.
10505239	10/12/04	TARASOVA ET AL.	229694

EXAMINER

Marcela M. Cordero Garcia

ART UNIT	PAPER
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1654 20071215

DATE MAILED:

Please find below and/or attached an Office communication concerning this application or proceeding.

Commissioner for Patents

This application contains sequence disclosures that are encompassed by the definitions for nucleotide and/or amino acid sequences set forth in 37 C.F.R. § 1.821(a)(1) and (a)(2). However, this application fails to comply with the requirements of 37 C.F.R. §§ 1.821-1.825 for the reason(s) set forth on the attached Notice To Comply With Requirements For Patent Applications Containing Nucleotide Sequence And/Or Amino Acid Sequence Disclosures. Applicant must comply with the requirements of the sequence rules (37 CFR 1.821 - 1.825) before the application can be examined under 35 U.S.C. §§ 131 and 132.

APPLICANT IS GIVEN ONE MONTH FROM THE DATE OF THIS LETTER WITHIN WHICH TO COMPLY WITH THE SEQUENCE RULES, 37 C.F.R. §§ 1.821-1.825. Failure to comply with these requirements will result in ABANDONMENT of the application under 37 C.F.R. § 1.821(g). Extensions of time may be obtained by filing a petition accompanied by the extension fee under the provisions of 37 C.F.R. § 1.136. In no case may an applicant extend the period for response beyond the six month statutory period. Direct the response to the undersigned. Applicant is requested to return a copy of the attached Notice to Comply with the response.

Please direct all replies to the United States Patent and Trademark Office via one (1) of the following:

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Alexandria, VA 22314

Any inquiry concerning this communication or earlier communications from the examiner should be directed to Examiner Marcela M Cordero Garcia whose telephone number is (571) 272-2939. If attempts to reach the Examiner by telephone are unsuccessful, the Examiner's supervisor, Cecilia Tsang, whose telephone number is (571) 272-0562.


Marcela M Cordero Garcia
Patent Examiner
Art Unit 1654


Cecilia J. Teardo
Patent Examiner
Art Unit 1654, Center 1600

Notice to ComplyApplication No.
10/505,239Applicant(s)
Tarasova et al.

Examiner

Art Unit
1654

M M Cordero Garcia

NOTICE TO COMPLY WITH REQUIREMENTS FOR PATENT APPLICATIONS CONTAINING NUCLEOTIDE SEQUENCE AND/OR AMINO ACID SEQUENCE DISCLOSURES

Applicant must file the items indicated below within the time period set the Office action to which the Notice is attached to avoid abandonment under 35 U.S.C. § 133 (extensions of time may be obtained under the provisions of 37 CFR 1.136(a)).

The nucleotide and/or amino acid sequence disclosure contained in this application does not comply with the requirements for such a disclosure as set forth in 37 C.F.R. 1.821 - 1.825 for the following reason(s):

- 1. This application clearly fails to comply with the requirements of 37 C.F.R. 1.821-1.825. Applicant's attention is directed to the final rulemaking notice published at 55 FR 18230 (May 1, 1990), and 1114 OG 29 (May 15, 1990). If the effective filing date is on or after July 1, 1998, see the final rulemaking notice published at 63 FR 29620 (June 1, 1998) and 1211 OG 82 (June 23, 1998).
- 2. This application does not contain, as a separate part of the disclosure on paper copy, a "Sequence Listing" as required by 37 C.F.R. 1.821(c).
- 3. A copy of the "Sequence Listing" in computer readable form has not been submitted as required by 37 C.F.R. 1.821(e).
- 4. A copy of the "Sequence Listing" in computer readable form has been submitted. However, the content of the computer readable form does not comply with the requirements of 37 C.F.R. 1.822 and/or 1.823, as indicated on the attached copy of the marked -up "Raw Sequence Listing."
- 5. The computer readable form that has been filed with this application has been found to be damaged and/or unreadable as indicated on the attached CRF Diskette Problem Report. A Substitute computer readable form must be submitted as required by 37 C.F.R. 1.825(d).
- 6. The paper copy of the "Sequence Listing" is not the same as the computer readable from of the "Sequence Listing" as required by 37 C.F.R. 1.821(e).
- 7. Other:

Applicant Must Provide:

- An initial or substitute computer readable form (CRF) copy of the "Sequence Listing".
- An initial or substitute paper copy of the "Sequence Listing", as well as an amendment directing its entry into the specification.
- A statement that the content of the paper and computer readable copies are the same and, where applicable, include no new matter, as required by 37 C.F.R. 1.821(e) or 1.821(f) or 1.821(g) or 1.825(b) or 1.825(d).

For questions regarding compliance to these requirements, please contact:

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Sequence Count Sheet	Application No. 10/505,239	Applicant(s) Tarasova et al.	
	Examiner MM Cordero Garcia	Art Unit 1654	

DATE OF COUNT

Mark only one space below

- (CRFN)** (CRF is unreadable; use CRF Diskette Problem Report)
- (CRFD)** (CRF does not comply; use Notice to Comply)
- (CRFR)** (CRF required but none submitted; use Notice to Comply)
- (bona fide)** (second or subsequent letter to applicant reporting bona fide attempt to comply; use Notice to Comply and send copy of RSL)
- (non bona fide)** (second or subsequent letter to applicant reporting non-bona fide attempt to comply; use Notice to Comply and send copy of RSL)

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Sequence Listing could not be accepted.

If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).

Reviewer: Anne Corrigan

Timestamp: Thu Aug 02 17:20:00 EDT 2007

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Reviewer Comments:

<210> 28

<211> 14

<212> PRT

<213> Artificial

<220>

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<220>

<221> misc_feature

<222> (12)..(12)

<223> Xaa can be any naturally occurring amino acid

<220>

<221> misc_feature

<222> (13)..(13)

<223> Xaa = at position 13 is norleucine

<400> 28

Phe Ala Leu Ala Glu Glu Ala Tyr Gly Trp Xaa Asp Phe

1

5

10

The above <222> (13)..(13) response is incorrect: "Xaa" is not at position 13, "Asp" is.

Application No: 10505239 Version No: 2.0

Input Set:

Output Set:

Started: 2007-07-30 18:17:35.343
Finished: 2007-07-30 18:17:36.501
Elapsed: 0 hr(s) 0 min(s) 1 sec(s) 158 ms
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Total Errors: 0
No. of SeqIDs Defined: 28
Actual SeqID Count: 28

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Output Set:

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Finished: 2007-07-30 18:17:36.501
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Total Warnings: 28
Total Errors: 0
No. of SeqIDs Defined: 28
Actual SeqID Count: 28

Error code	Error Description
	This error has occurred more than 20 times, will not be displayed.

SEQUENCE LISTING

<110> TARASOVA, Nadya I
MICHEJDA, Christopher J
DYBA, Marcin
COHRAN, Carolyn

<120> CONJUGATES OF LIGAND, LINKER AND CYTOTOXIC AGENT AND RELATED
COMPOSITIONS AND METHODS OF USE

<130> 229694

<140> 10505239
<141> 2004-10-12

<150> US 10/505,239
<151> 2004-10-12

<150> PCT/US03/06344
<151> 2003-02-27

<150> 60/360,543
<151> 2002-02-27

<150> 60/370,189
<151> 2002-04-05

<160> 28

<170> PatentIn version 3.4

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Gln Gly Pro Trp Leu Glu Glu Glu Ala Tyr Gly Trp Met Asp
20 25 30

Phe

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1 5

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1 5 10 15

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20 25

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<213> Artificial Sequence

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Trp Ala Val Gly His Leu Met

1 5

<210> 11

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<212> PRT

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<210> 12

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<223> wherein the peptide is carboxylated at either the N-or C-terminus

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Arg Pro Leu Pro Gln Gln Phe Phe Gly Leu Met

1 5 10

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<211> 15

<212> PRT

<213> Artificial Sequence

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1 5 10 15

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1

5

10

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Met Ala Val Lys Lys Tyr Leu Asn Ser Ile Leu Asn Gly
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Xaa Ala Val Lys Lys Tyr Leu Asn Ser Ile Leu Asn Gly
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Xaa Xaa Leu Ala Leu

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Xaa Xaa Leu Ala Leu

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1 5

<210> 25
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<223> F = at position 15 comprises a C-terminal amide group

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Xaa Leu Ala Leu Ala Glu Glu Ala Tyr Gly Trp Xaa Asp Phe
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<210> 28
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<220>

<221> misc_feature
<222> (13)..(13)
<223> Xaa = at position 13 is norleucine

<400> 28

Phe Ala Leu Ala Glu Glu Ala Tyr Gly Trp Xaa Asp Phe
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